



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/019,495

DATE: 06/04/2002

TIME: 14:51:10

Input Set : A:\pf0698usn.txt

Output Set: N:\CRF3\06042002\J019495.raw

1 <110> APPLICANT: Hillman, Jennifer L.
 2 Bandman, Olga
 3 Tang, Y. Tom
 4 Lal, Preeti
 5 Yue, Henry
 6 Reddy, Roopa
 7 Azimzai, Yalda
 8 Baughn, Mariah R.
 10 <120> TITLE OF INVENTION: HUMAN MEMBRANE-ASSOCIATED PROTEINS
 12 <130> FILE REFERENCE: PF-0698 PCT
 C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/019,495
 C--> 15 <141> CURRENT FILING DATE: 2002-04-15
 17 <160> NUMBER OF SEQ ID NOS: 34
 19 <170> SOFTWARE: PERL Program
 23 <210> SEQ ID NO: 1
 24 <211> LENGTH: 175
 25 <212> TYPE: PRT
 26 <213> ORGANISM: Homo sapiens
 28 <220> FEATURE:
 29 <223> OTHER INFORMATION: 402771CD1
 31 <400> SEQUENCE: 1
 32 Met Ser His Gly Ala Gly Leu Val Arg Thr Thr Cys Ser Ser Gly
 33 1 5 10 15
 34 Ser Ala Leu Gly Pro Gly Ala Gly Ala Gln Pro Ser Ala Ser
 35 20 25 30
 36 Pro Leu Glu Gly Leu Leu Asp Leu Ser Tyr Pro Arg Thr His Ala
 37 35 40 45
 38 Ala Leu Leu Lys Val Ala Gln Met Val Thr Leu Leu Ile Ala Phe
 39 50 55 60
 40 Ile Cys Val Arg Ser Ser Leu Trp Thr Asn Tyr Ser Ala Tyr Ser
 41 65 70 75
 42 Tyr Phe Glu Val Val Thr Ile Cys Asp Leu Ile Met Ile Leu Ala
 43 80 85 90
 44 Phe Tyr Leu Val His Leu Phe Arg Phe Tyr Arg Val Leu Thr Cys
 45 95 100 105
 46 Ile Ser Trp Pro Leu Ser Glu Leu Leu His Tyr Leu Ile Gly Thr
 47 110 115 120
 48 Leu Leu Leu Leu Ile Ala Ser Ile Val Ala Ala Ser Lys Ser Tyr
 49 125 130 135
 50 Asn Gln Ser Gly Leu Val Ala Gly Ala Ile Phe Gly Phe Met Ala
 51 140 145 150
 52 Thr Phe Leu Cys Met Ala Ser Ile Trp Leu Ser Tyr Lys Ile Ser
 53 155 160 165

ENTERED

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54 Cys Val Thr Gln Ser Thr Asp Ala Ala Val
55           170           175
58 <210> SEQ ID NO: 2
59 <211> LENGTH: 161
60 <212> TYPE: PRT
61 <213> ORGANISM: Homo sapiens
63 <220> FEATURE:
64 <223> OTHER INFORMATION: 1296216CD1
66 <400> SEQUENCE: 2
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68   1           5           10           15
69 Thr Ser Glu Thr Gly Arg Ala Leu Thr Leu Leu Ala Gln Gln Glu
70           20           25           30
71 Thr Gln Met Arg Asn Ala Ile Tyr Gln Asn Arg Leu Ala Leu Asp
72           35           40           45
73 Tyr Leu Leu Ala Ala Glu Gly Gly Val Cys Glu Lys Phe Asn Leu
74           50           55           60
75 Thr Lys Cys Cys Leu Gln Arg Asp Asp Gln Gly Gln Val Val Lys
76           65           70           75
77 Asn Ile Val Arg Asp Met Thr Lys Leu Ala His Val Pro Met Gln
78           80           85           90
79 Val Trp His Arg Phe Asp Pro Gly Ser Leu Phe Gly Lys Trp Leu
80           95          100          105
81 Pro Ala Leu Gly Gly Phe Lys Thr Leu Ile Ile Gly Met Ile Met
82          110          115          120
83 Val Leu Gly Thr Cys Met Leu Leu Pro Cys Met Leu Pro Ile Phe
84          125          130          135
85 Leu Gln Leu Leu Arg Ser Phe Val Ile Thr Leu Val His Gln Lys
86          140          145          150
87 Thr Ser Ala Gln Val Tyr Tyr Met Asn His Tyr
88          155          160
91 <210> SEQ ID NO: 3
92 <211> LENGTH: 563
93 <212> TYPE: PRT
94 <213> ORGANISM: Homo sapiens
96 <220> FEATURE:
97 <223> OTHER INFORMATION: 1693173CD1
99 <400> SEQUENCE: 3
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101   1           5           10           15
102 Leu Phe His Phe Gly Met Thr Trp Gln Ile Asn Gly Thr Gly Leu
103           20           25           30
104 Leu Gly Lys Arg Asn Asp Gln Leu Pro Ser Asn Phe Thr Pro Val
105           35           40           45
106 Phe Tyr Ser Gln Leu Gln Lys Asn Ser Ser Trp Ala Glu His Leu
107           50           55           60
108 Ile Ser Asn Cys Asp Gly Asp Ser Ser Cys Ile Tyr Asp Thr Leu
109           65           70           75
110 Ala Leu Arg Asn Ala Ser Ile Gly Leu His Thr Arg Glu Val Ser

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111		80		85		90
112	Lys Asn Tyr Glu Gln Ala Asn Ala Thr Leu Asn Gln Tyr Pro Pro					
113		95		100		105
114	Ser Ile Asn Gly Gly Arg Val Ile Glu Ala Tyr Lys Gly Gln Thr					
115		110		115		120
116	Thr Leu Ile Gln Tyr Thr Ser Asn Ala Glu Asp Ala Asn Phe Thr					
117		125		130		135
118	Leu Arg Asp Ser Cys Thr Asp Leu Glu Leu Phe Glu Asn Gly Thr					
119		140		145		150
120	Leu Leu Trp Thr Pro Lys Ser Leu Glu Pro Phe Thr Leu Glu Ile					
121		155		160		165
122	Leu Ala Arg Ser Ala Lys Ile Gly Leu Ala Ser Ala Leu Gln Pro					
123		170		175		180
124	Arg Thr Val Val Cys His Cys Asn Ala Glu Ser Gln Cys Leu Tyr					
125		185		190		195
126	Asn Gln Thr Ser Arg Val Gly Asn Ser Ser Leu Glu Val Ala Gly					
127		200		205		210
128	Cys Lys Cys Asp Gly Gly Thr Phe Gly Arg Tyr Cys Glu Gly Ser					
129		215		220		225
130	Glu Asp Ala Cys Glu Glu Pro Cys Phe Pro Ser Val His Cys Val					
131		230		235		240
132	Pro Gly Lys Gly Cys Glu Ala Cys Pro Pro Asn Leu Thr Gly Asp					
133		245		250		255
134	Gly Arg His Cys Ala Ala Leu Gly Ser Ser Phe Leu Cys Gln Asn					
135		260		265		270
136	Gln Ser Cys Pro Val Asn Tyr Cys Tyr Asn Gln Gly His Cys Tyr					
137		275		280		285
138	Ile Ser Gln Thr Leu Gly Cys Gln Pro Met Cys Thr Cys Pro Pro					
139		290		295		300
140	Ala Phe Thr Asp Ser Arg Cys Phe Leu Ala Gly Asn Asn Phe Ser					
141		305		310		315
142	Pro Thr Val Asn Leu Glu Leu Pro Leu Arg Val Ile Gln Leu Leu					
143		320		325		330
144	Leu Ser Glu Glu Glu Asn Ala Ser Met Ala Glu Val Asn Ala Ser					
145		335		340		345
146	Val Ala Tyr Arg Leu Gly Thr Leu Asp Met Arg Ala Phe Leu Arg					
147		350		355		360
148	Asn Ser Gln Val Glu Arg Ile Asp Ser Ala Ala Pro Ala Ser Gly					
149		365		370		375
150	Ser Pro Ile Gln His Trp Met Val Ile Ser Glu Phe Gln Tyr Arg					
151		380		385		390
152	Pro Arg Gly Pro Val Ile Asp Phe Leu Asn Asn Gln Leu Leu Ala					
153		395		400		405
154	Ala Val Val Glu Ala Phe Leu Tyr His Val Pro Arg Arg Ser Glu					
155		410		415		420
156	Glu Pro Arg Asn Asp Val Val Phe Gln Pro Ile Ser Gly Glu Asp					
157		425		430		435
158	Val Arg Asp Val Thr Ala Leu Asn Val Ser Thr Leu Lys Ala Tyr					
159		440		445		450

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```

160 Phe Arg Cys Asp Gly Tyr Lys Gly Tyr Asp Leu Val Tyr Ser Pro
161      455      460      465
162 Gln Ser Gly Phe Thr Cys Val Ser Pro Cys Ser Arg Gly Tyr Cys
163      470      475      480
164 Asp His Gly Gly Gln Cys Gln His Leu Pro Ser Gly Pro Arg Cys
165      485      490      495
166 Ser Cys Val Ser Phe Ser Ile Tyr Thr Ala Trp Gly Glu His Cys
167      500      505      510
168 Glu His Leu Ser Met Lys Leu Asp Ala Phe Phe Gly Ile Phe Phe
169      515      520      525
170 Gly Val Leu Gly Gly Leu Leu Leu Leu Gly Val Gly Thr Phe Val
171      530      535      540
172 Val Leu Arg Phe Trp Gly Cys Ser Gly Ala Arg Phe Ser Tyr Phe
173      545      550      555
174 Leu Asn Ser Ala Glu Ala Leu Pro
175      560
178 <210> SEQ ID NO: 4
179 <211> LENGTH: 396
180 <212> TYPE: PRT
181 <213> ORGANISM: Homo sapiens
183 <220> FEATURE:
184 <223> OTHER INFORMATION: 2095069CD1
186 <400> SEQUENCE: 4
187 Met Ala Val Gln Glu Ser Ala Ala Gln Leu Ser Met Thr Leu Lys
188   1      5      10      15
189 Val Gln Glu Tyr Pro Thr Leu Lys Val Pro Tyr Glu Thr Leu Asn
190      20      25      30
191 Lys Arg Phe Arg Ala Ala Gln Lys Asn Ile Asp Arg Glu Thr Ser
192      35      40      45
193 His Val Thr Met Val Val Ala Glu Leu Glu Lys Thr Leu Ser Gly
194      50      55      60
195 Cys Pro Ala Val Asp Ser Val Val Ser Leu Leu Asp Gly Val Val
196      65      70      75
197 Glu Lys Leu Ser Val Leu Lys Arg Lys Ala Val Glu Ser Ile Gln
198      80      85      90
199 Ala Glu Asp Glu Ser Ala Lys Leu Cys Lys Arg Arg Ile Glu His
200      95     100     105
201 Leu Lys Glu His Ser Ser Asp Gln Pro Ala Ala Ala Ser Val Trp
202     110     115     120
203 Lys Arg Lys Arg Met Asp Arg Met Met Val Glu His Leu Leu Arg
204     125     130     135
205 Cys Gly Tyr Tyr Asn Thr Ala Val Lys Leu Ala Arg Gln Ser Gly
206     140     145     150
207 Ile Glu Asp Leu Val Asn Ile Glu Met Phe Leu Thr Ala Lys Glu
208     155     160     165
209 Val Glu Glu Ser Leu Glu Arg Arg Glu Thr Ala Thr Cys Leu Ala
210     170     175     180
211 Trp Cys His Asp Asn Lys Ser Arg Leu Arg Lys Met Lys Ser Cys
212     185     190     195

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```

213 Leu Glu Phe Ser Leu Arg Ile Gln Glu Phe Ile Glu Leu Ile Arg
214          200          205          210
215 Gln Asn Lys Arg Leu Asp Ala Val Arg His Ala Arg Lys His Phe
216          215          220          225
217 Ser Gln Ala Glu Gly Ser Gln Leu Asp Glu Val Arg Gln Ala Met
218          230          235          240
219 Gly Met Leu Ala Phe Pro Pro Asp Thr His Ile Ser Pro Tyr Lys
220          245          250          255
221 Asp Leu Leu Asp Pro Ala Arg Trp Arg Met Leu Ile Gln Gln Phe
222          260          265          270
223 Arg Tyr Asp Asn Tyr Arg Leu His Gln Leu Gly Asn Asn Ser Val
224          275          280          285
225 Phe Thr Leu Thr Leu Gln Ala Gly Leu Ser Ala Ile Lys Thr Pro
226          290          295          300
227 Gln Cys Tyr Lys Glu Asp Gly Ser Ser Lys Ser Pro Asp Cys Pro
228          305          310          315
229 Val Cys Ser Arg Ser Leu Asn Lys Leu Ala Gln Pro Leu Pro Met
230          320          325          330
231 Ala His Cys Ala Asn Ser Arg Leu Val Cys Lys Ile Ser Gly Asp
232          335          340          345
233 Val Met Asn Glu Asn Asn Pro Pro Met Met Leu Pro Asn Gly Tyr
234          350          355          360
235 Val Tyr Gly Tyr Asn Ser Leu Leu Ser Ile Arg Gln Asp Asp Lys
236          365          370          375
237 Val Val Cys Pro Arg Thr Lys Glu Val Phe His Phe Ser Gln Ala
238          380          385          390
239 Glu Lys Val Tyr Ile Met
240          395
243 <210> SEQ ID NO: 5
244 <211> LENGTH: 265
245 <212> TYPE: PRT
246 <213> ORGANISM: Homo sapiens
248 <220> FEATURE:
249 <223> OTHER INFORMATION: 2645927CD1
251 <400> SEQUENCE: 5
252 Met Asn Met Ser Val Leu Thr Leu Gln Glu Tyr Glu Phe Glu Lys
253   1          5          10          15
254 Gln Phe Asn Glu Asn Glu Ala Ile Gln Trp Met Gln Glu Asn Trp
255          20          25          30
256 Lys Lys Ser Phe Leu Phe Ser Ala Leu Tyr Ala Ala Phe Ile Phe
257          35          40          45
258 Gly Gly Arg His Leu Met Asn Lys Arg Ala Lys Phe Glu Leu Arg
259          50          55          60
260 Lys Pro Leu Val Leu Trp Ser Leu Thr Leu Ala Val Phe Ser Ile
261          65          70          75
262 Phe Gly Ala Leu Arg Thr Gly Ala Tyr Met Val Tyr Ile Leu Met
263          80          85          90
264 Thr Lys Gly Leu Lys Gln Ser Val Cys Asp Gln Gly Phe Tyr Asn
265          95          100          105

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RAW SEQUENCE LISTING ERROR SUMMARY
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:31; N Pos. 3607